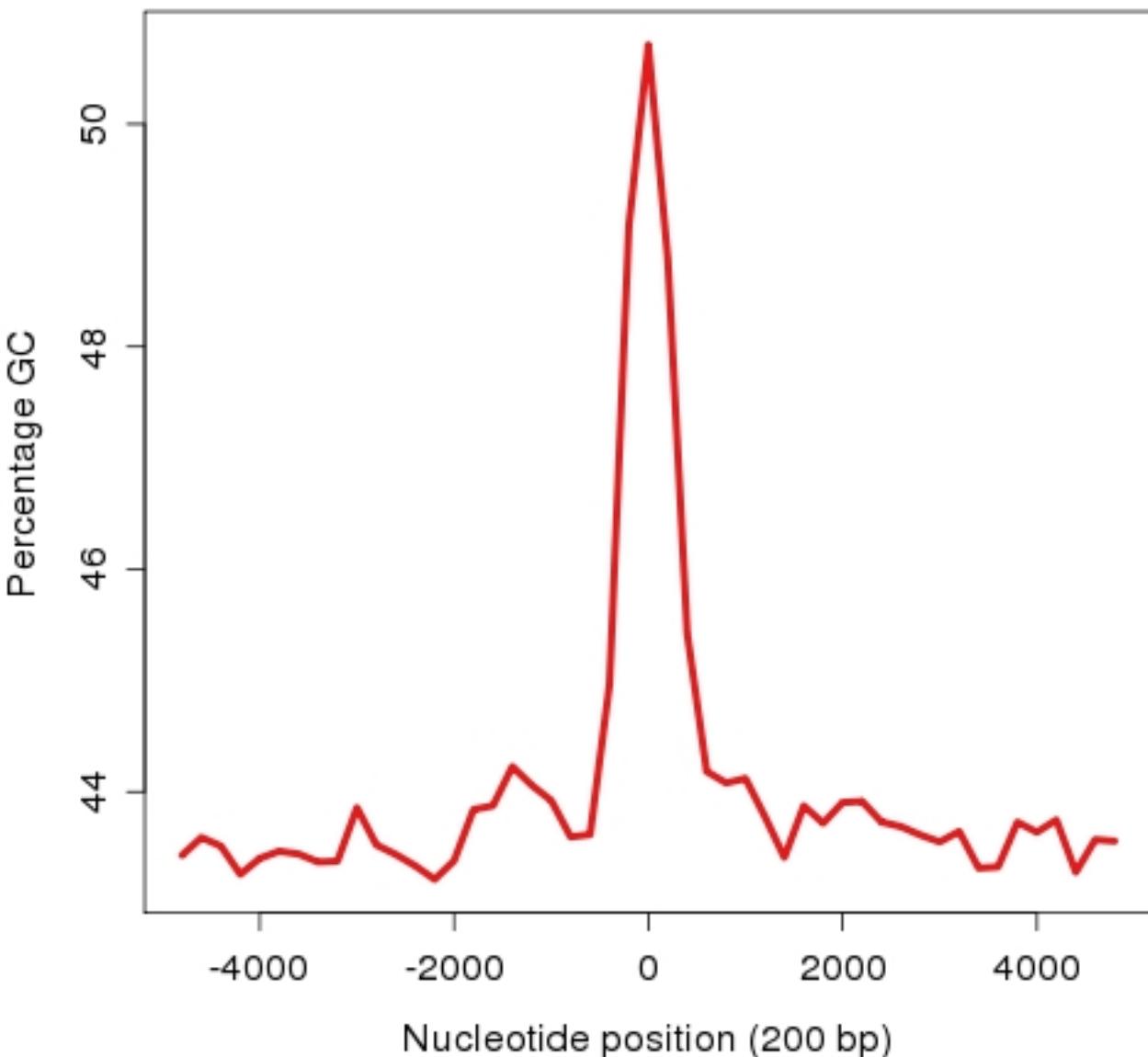
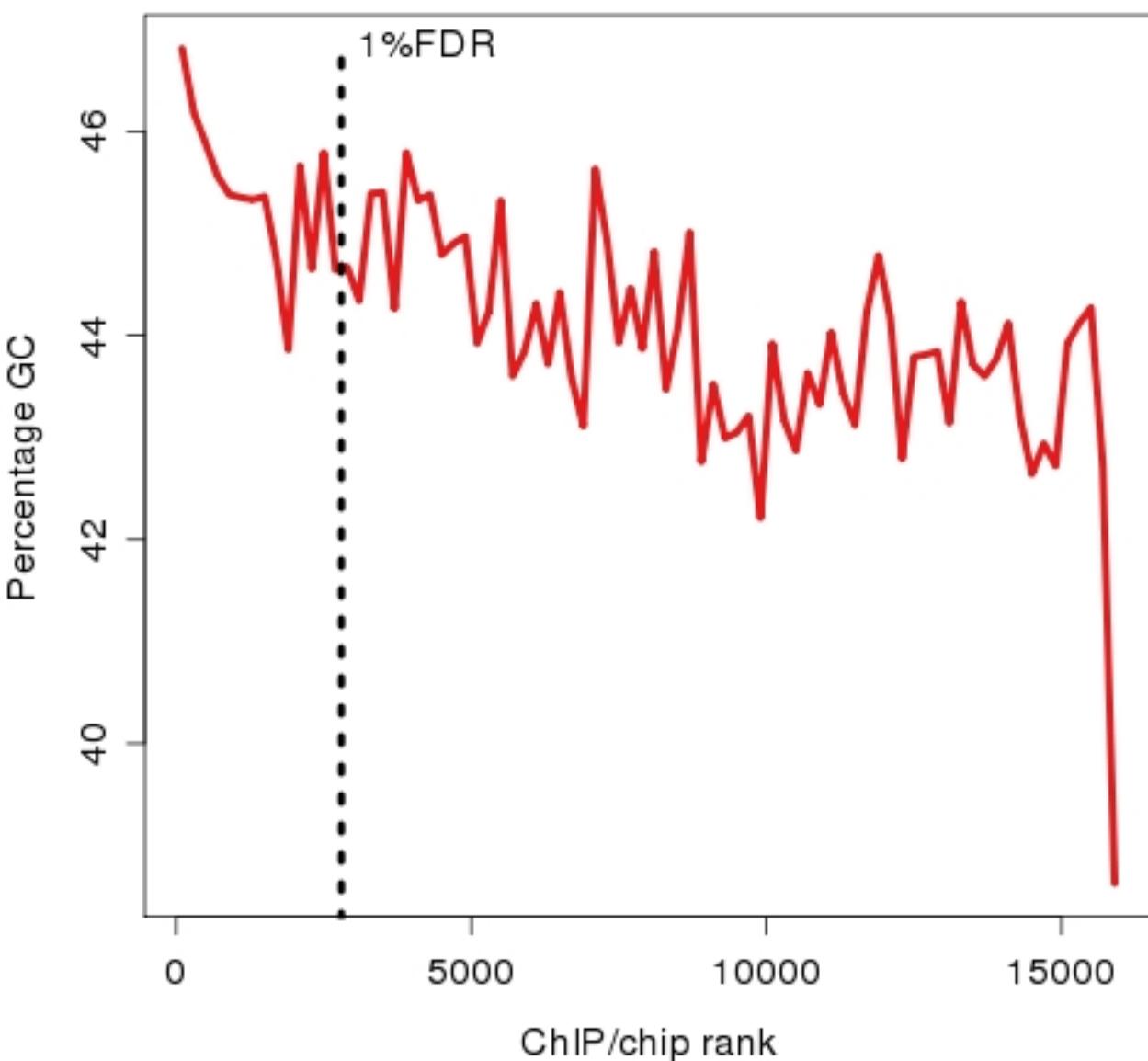


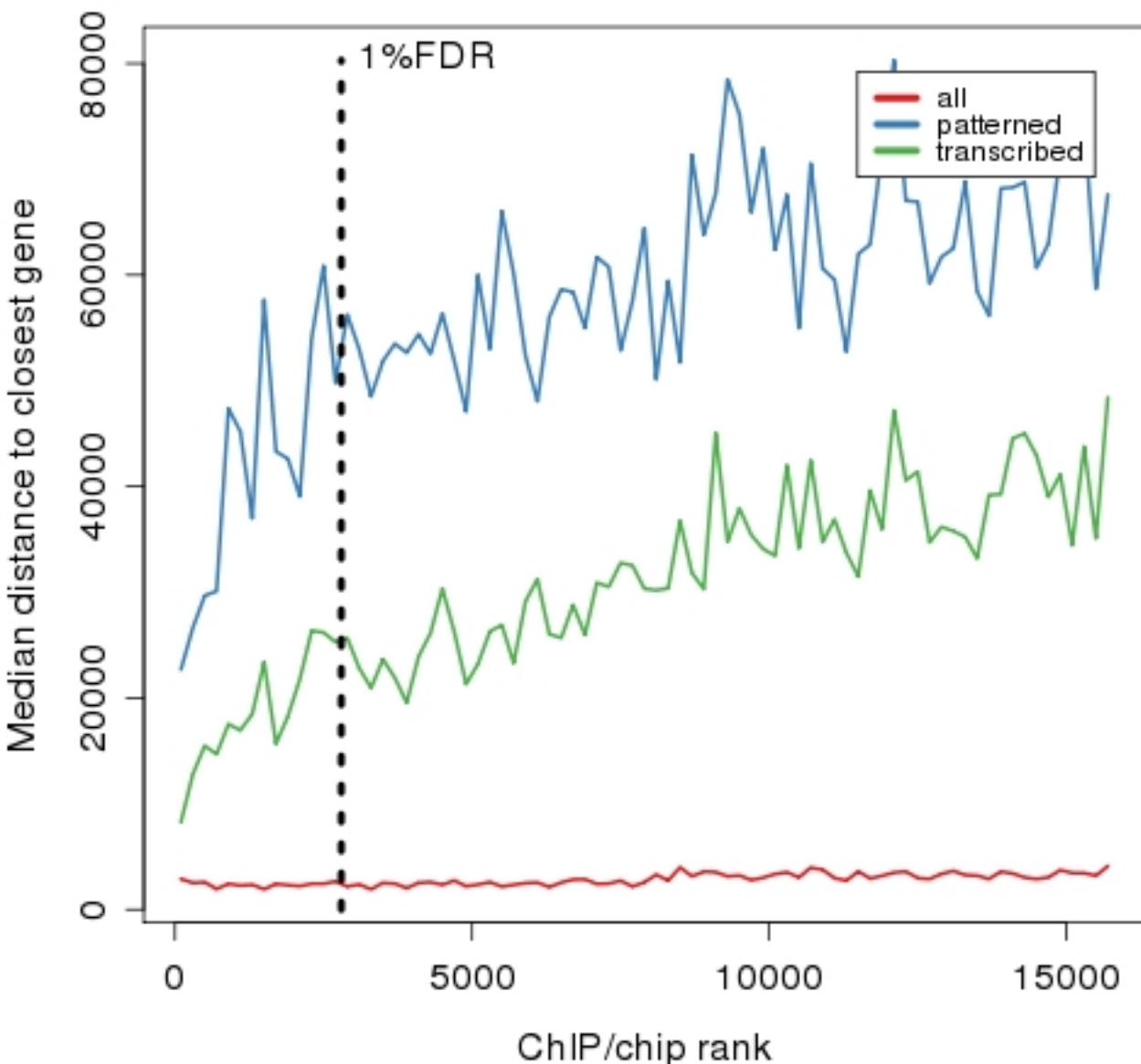
## **sna\_2 base composition across peaks**



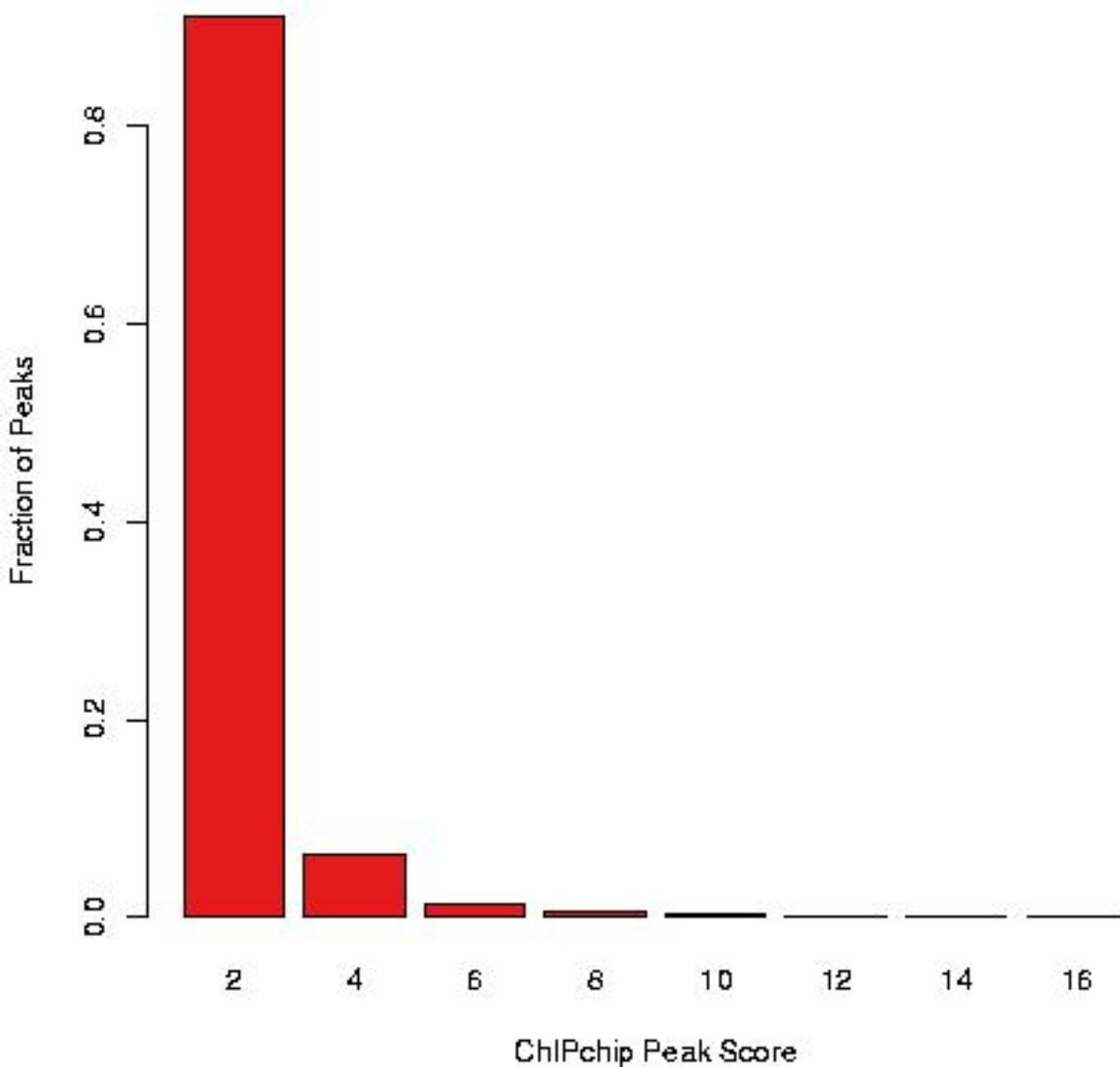
## sna\_2 base composition down ranks



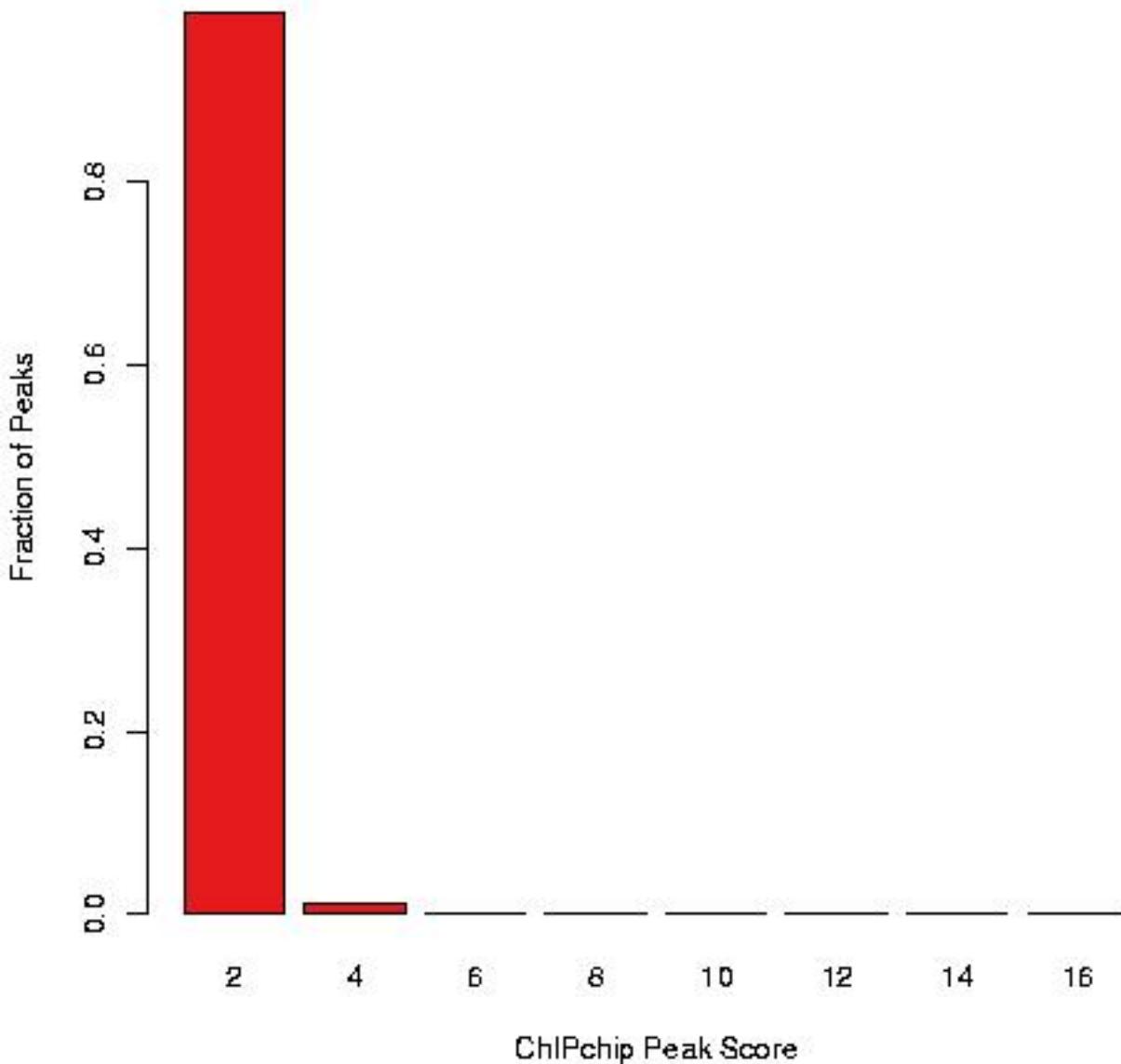
## sna\_2 median distance to genes



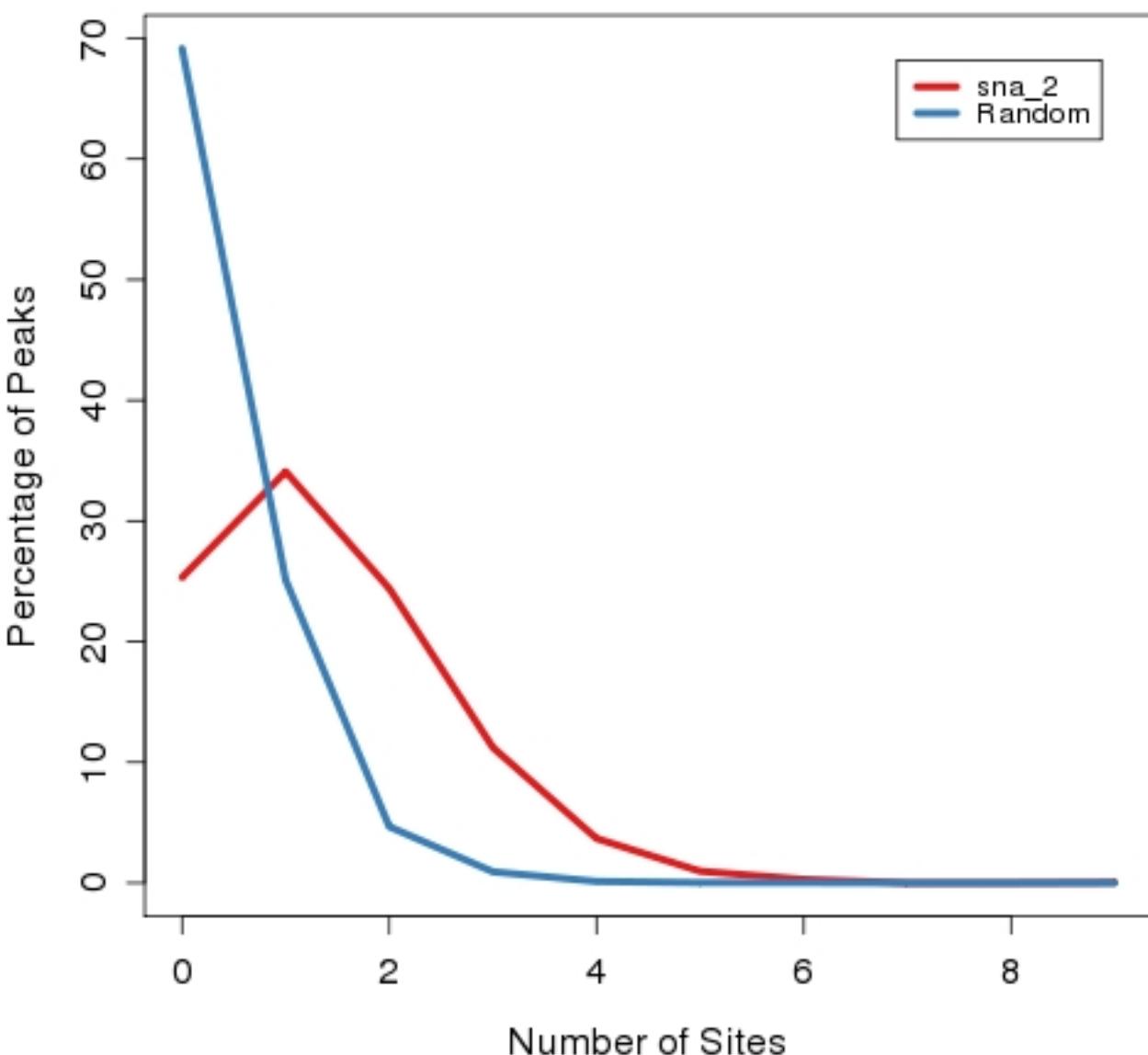
### sna\_2 distribution of 1%FDR primary peak scores



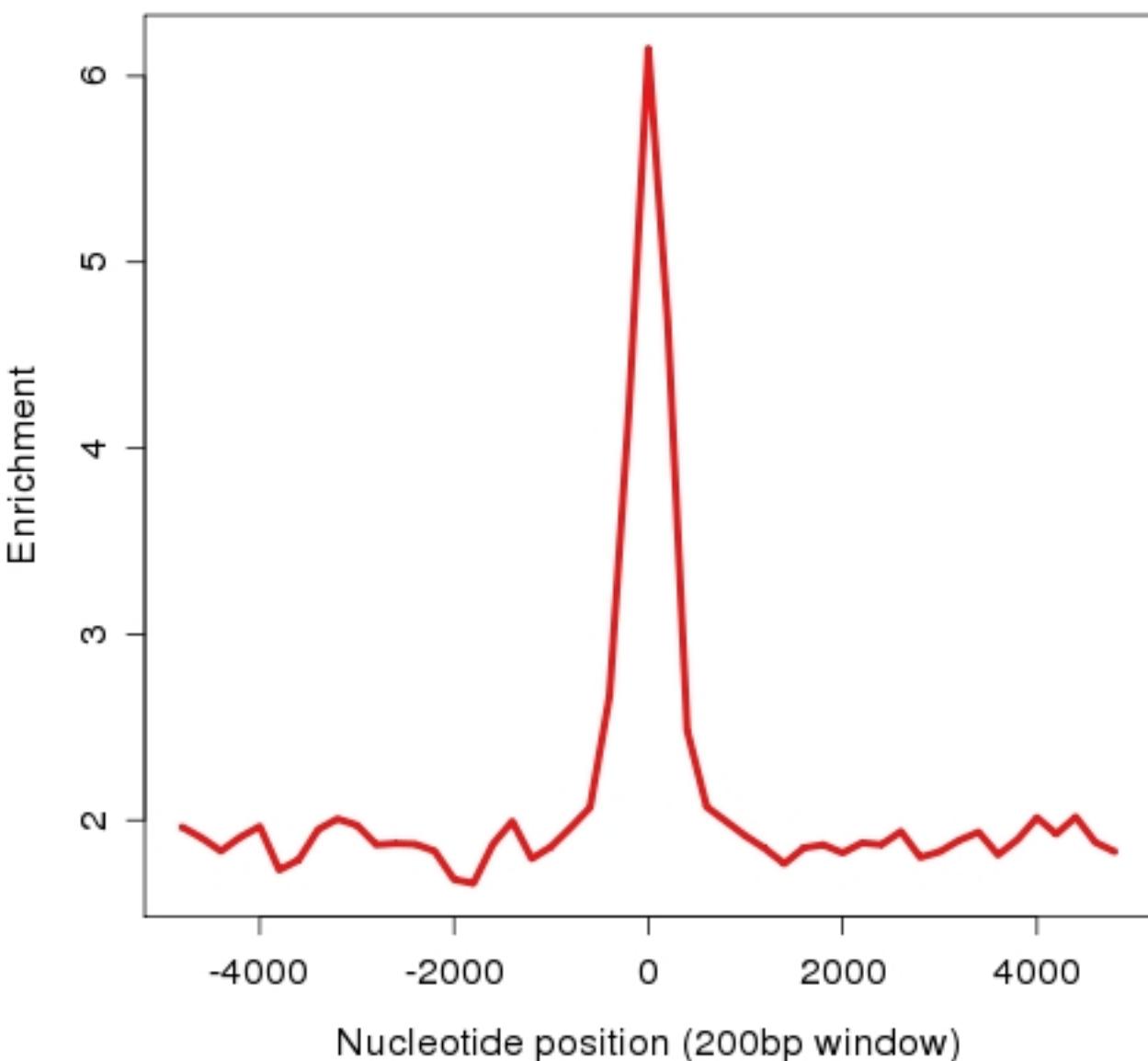
### sna\_2 distribution of 25%FDR primary peak scores



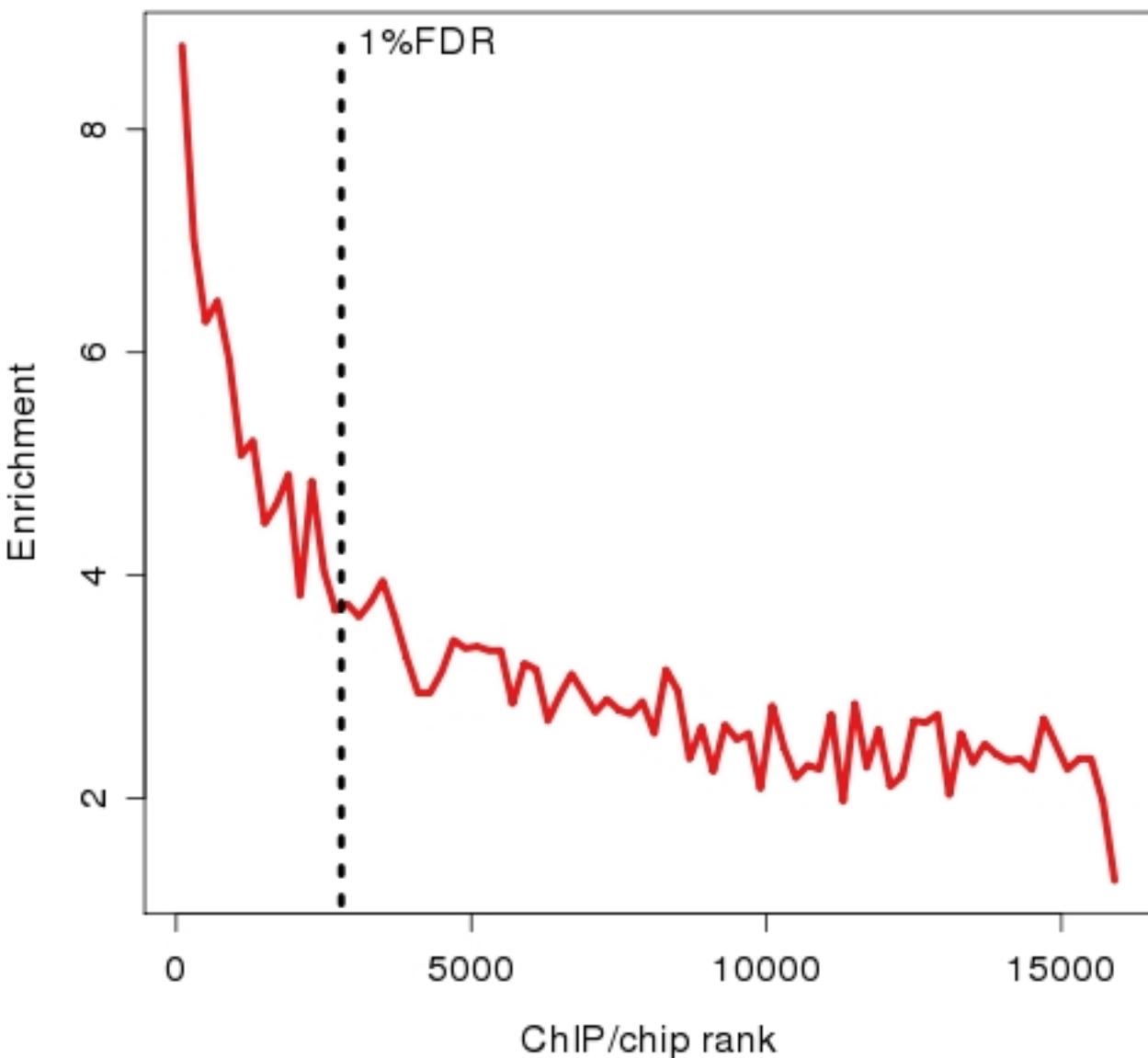
## **sna\_2 distribution of number of sites**



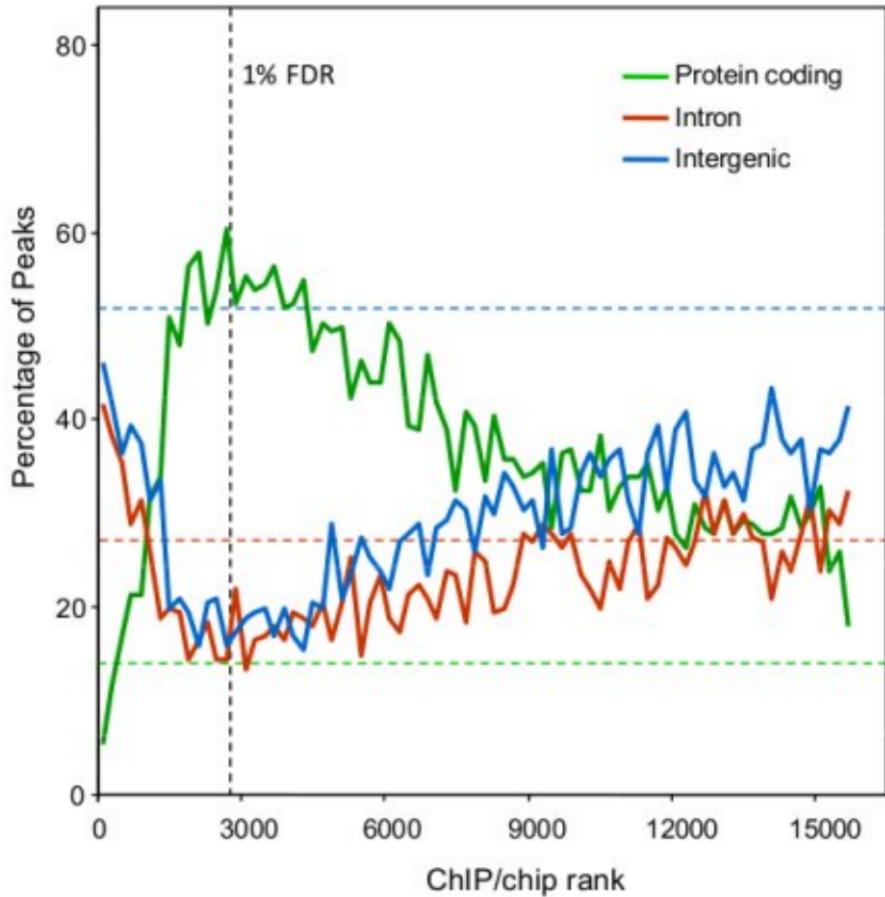
## sna\_2 PWM enrichment across peaks



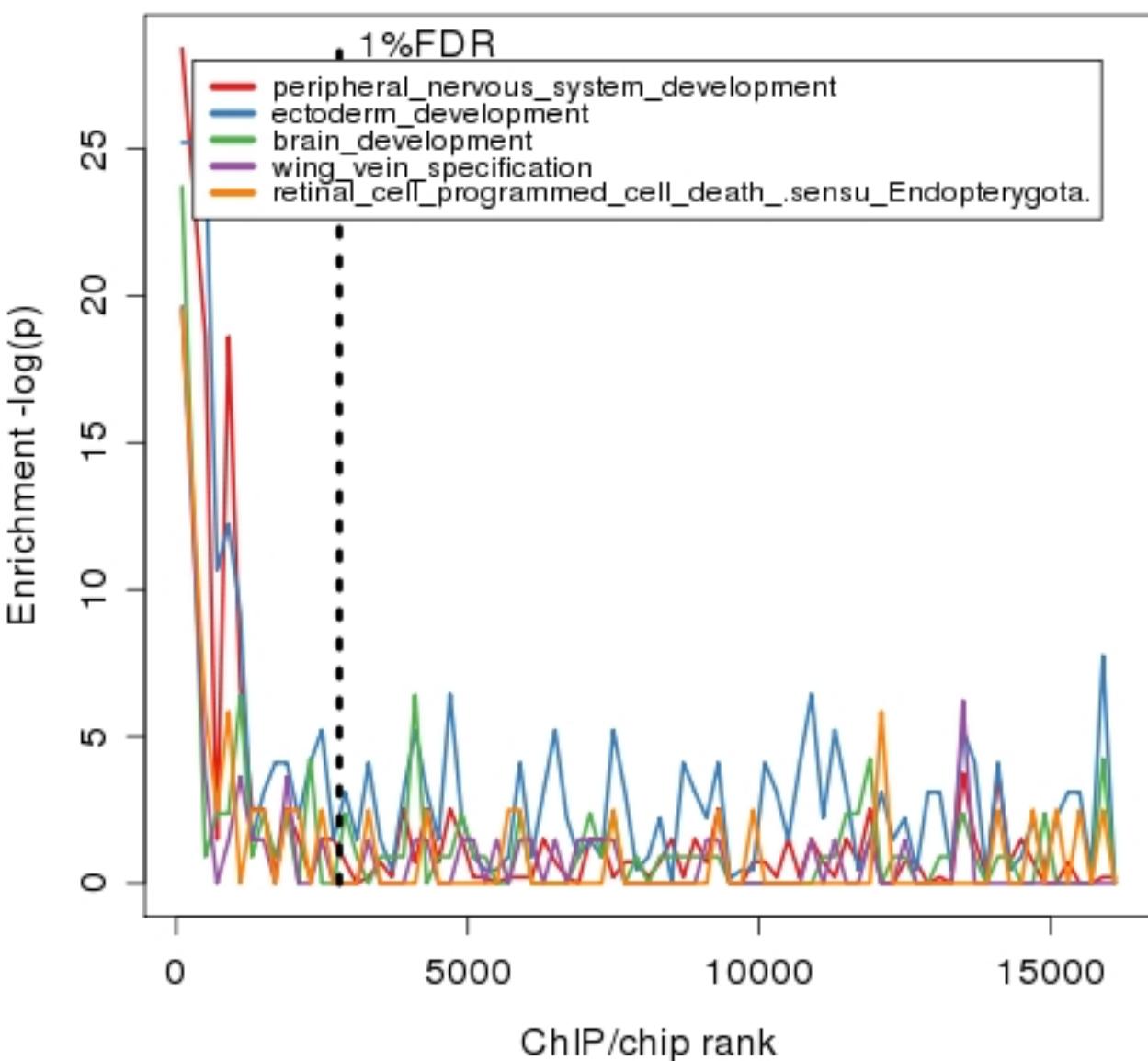
## sna\_2 PWM enrichment down ranks

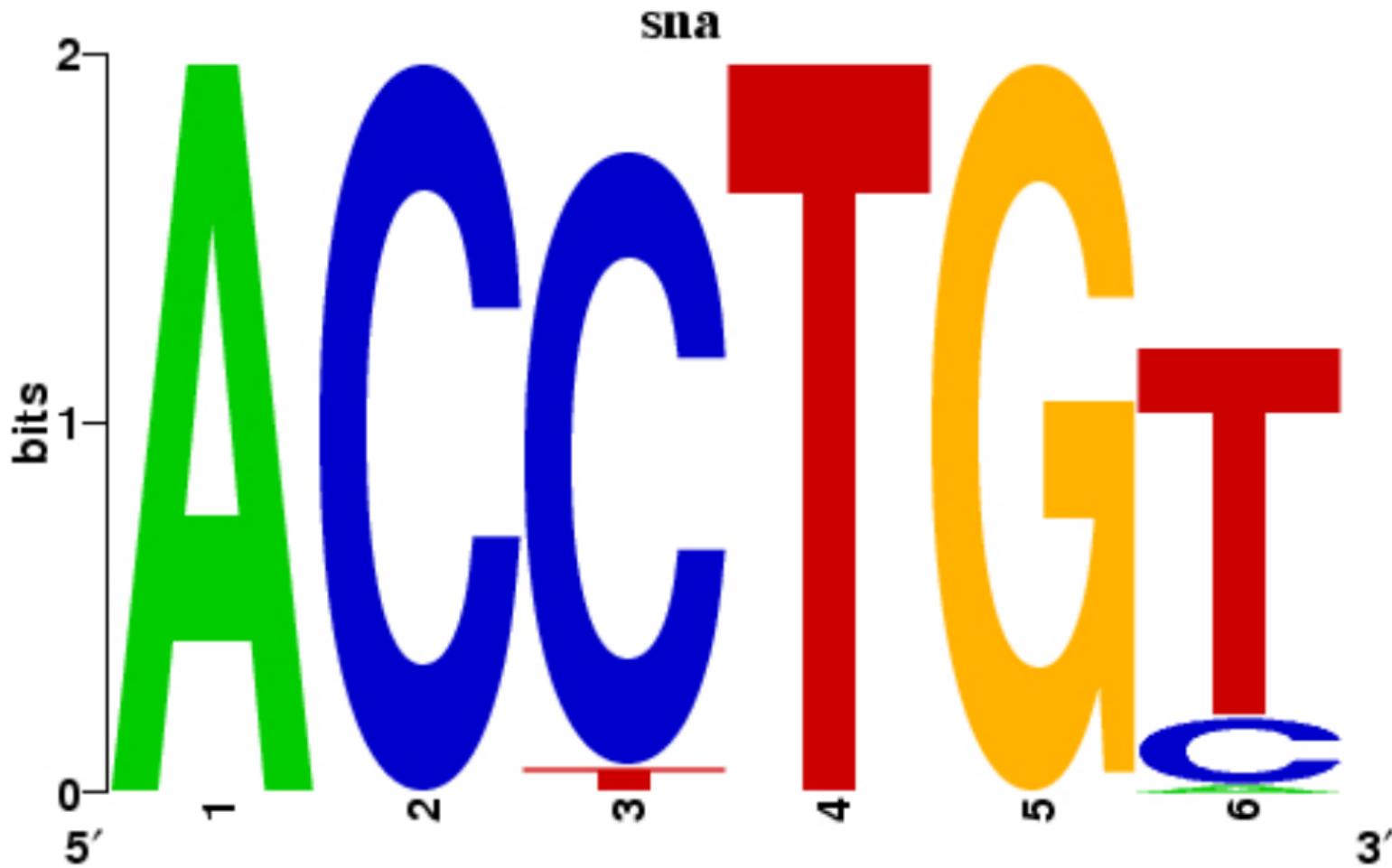


## SNA 2 genomic location of peaks

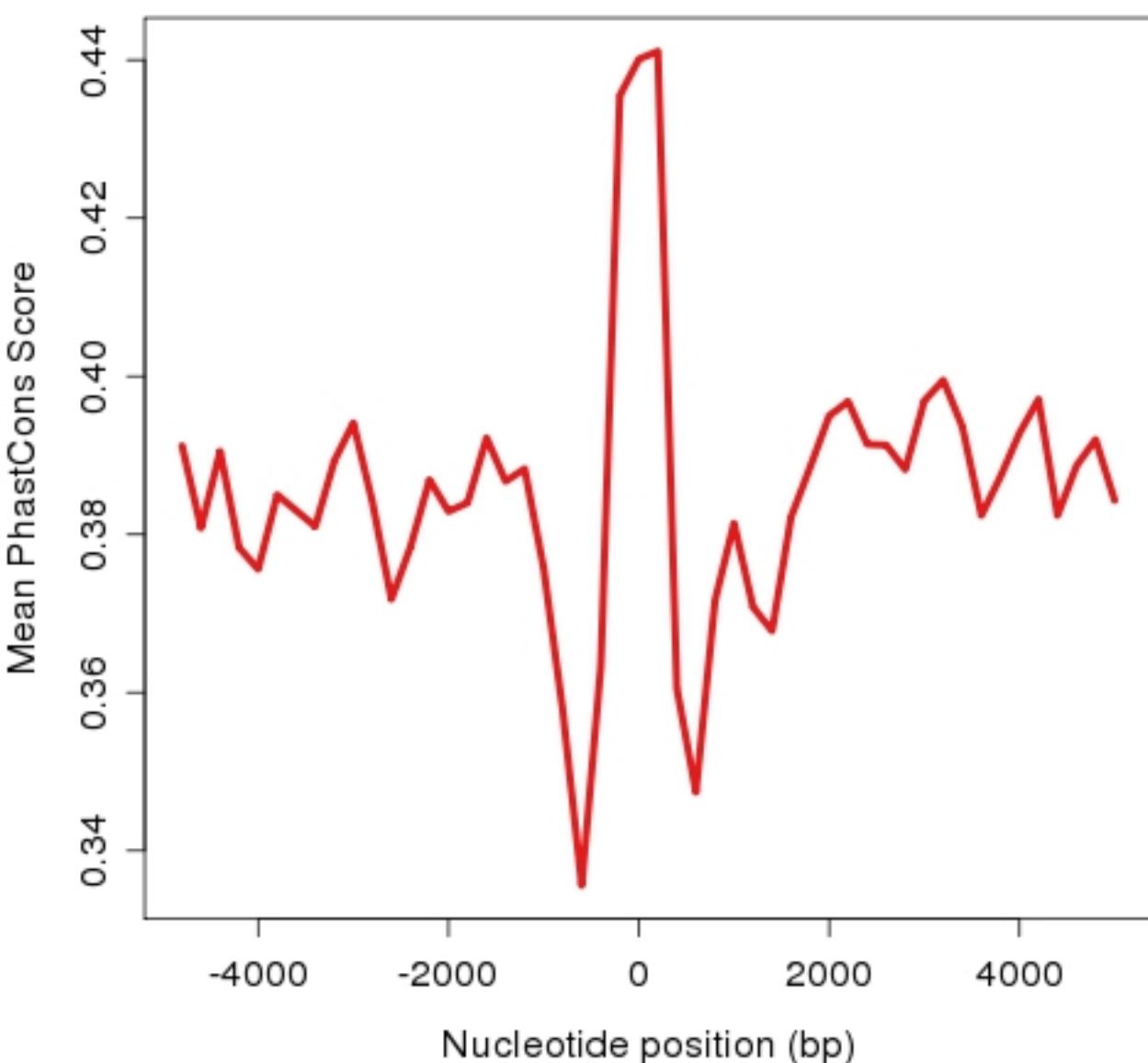


## sna\_2 GO term enrichment (-logP)

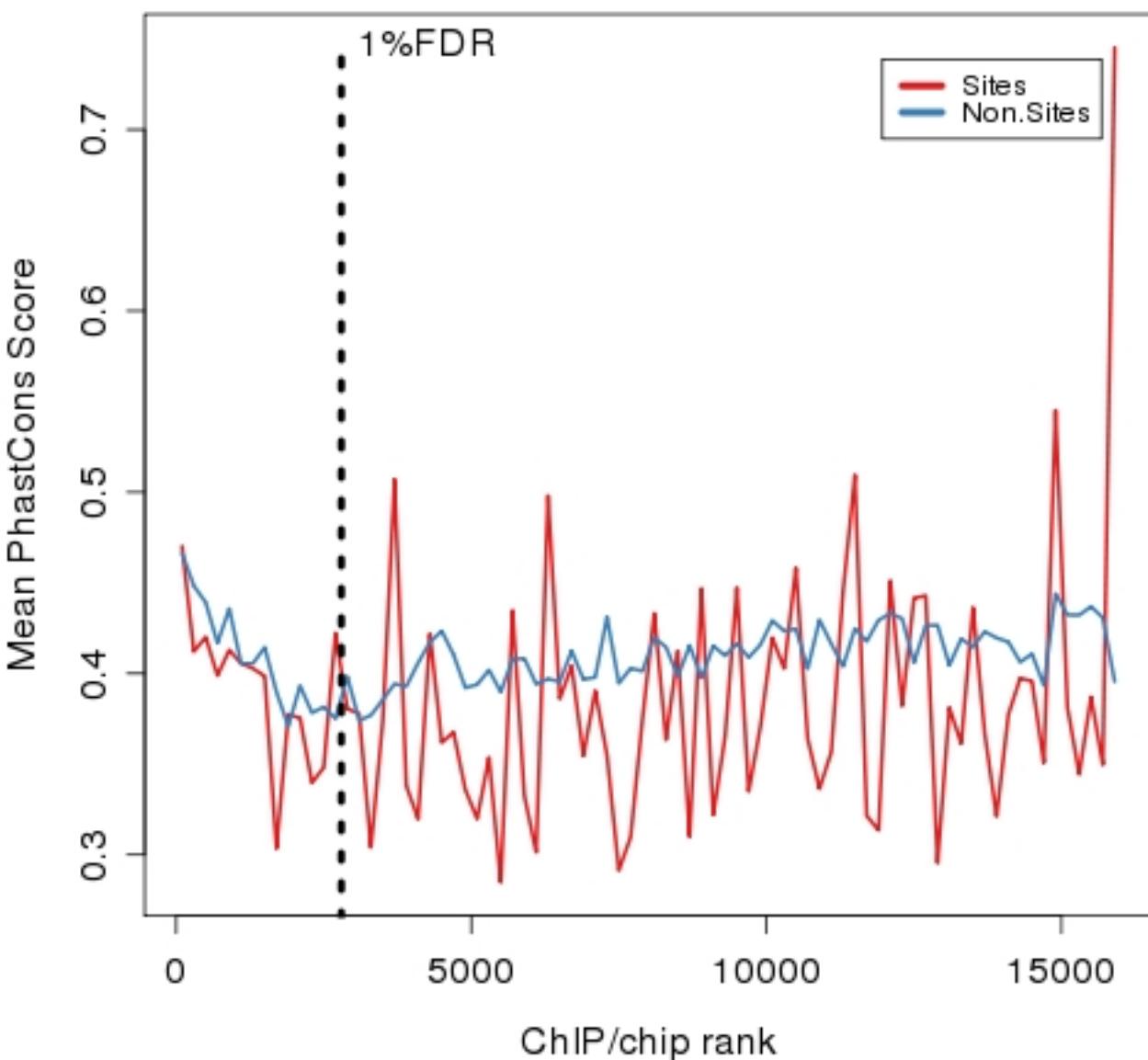




## sna\_2 mean PhastCons across peaks



## sna\_2 PhastCons



## sna\_2 mean PhastCons down ranks

